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954.2
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length: 2000000000
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A. faecalis nitril
Plasmid pKK232-8
Plasmid pCM-B64
Plasmid pCM-P1208
Plasmid pCM-P1201
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ALIGNMENTS

RESULT AAA47190

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AAA47190 standard; DNA; 1793

ВP

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CDS
                                                                                                                                             Synthetic.
                                                                                                                                                            nitrile hydratase; amidase; pRPA-BCAT41;
                                                                                                                                                                                     Nucleotide sequence of the expression cassette of pRPA-BCAT41.
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                                                                                                                                                                                                                     AAA47190;
       Favre-Bulle 0,
                      (RHON ) RHONE-POULENC ANIMAL NUTRITION SA
                                       11-DEC-1998;
19-JUL-1999;
                                                              10-DEC-1999;
                                                                              22-JUN-2000
                                                                                              WO200036120-A1
                                                                                                                                                                     Methionine; 2-hydroxy-4-methylthiobutanoic acid; nitrilase;
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                                       98FR-0015849
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Best Local Similarity 100
Matches 1793; Conservative
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CC This sequence encodes the nitrilase enzyme from Alcaligenes faecalis. CC The invention relates to a method of preparing organic acid by hydrolysing nitrile compounds of formula RICH(R2)CN with nitrilase to CC form carboxylic acid of formula RICH(R2)COOH and equimolar amount of CC ammonia. (R1 is optionally substituted phenyl, naphthyl, indole, CC thiophene, pyridyl or cyclohexenyl; R1 is H or hydroxyl group). The CC nitrilase is prepared by culturing a transformant containing recombinant CC DNA obtained by integrating into a vector the A. faecalis nitrilase CC enzyme DNA. Various kinds of amide or carboxylic acid can be produced CC industrially in good efficiency.

XX Sequence 1727 BP; 343 A; 566 C; 468 G; 350 T; 0 other;
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                                                                                                     This invention describes a novel nucleic acid sequence (I) encody Alcaligenes faecalis nitrilase polypeptide. The polypeptide encody (I), or a growing, restling or digested microorganism containing useful for preparing chiral carboxylic acids (III) by conversion racemic nitriles (IV). (III) are useful as intermediates, e.g. for pharmaceuticals and plant protection agents. This sequence encode Alcaligenes faecalis nitrilase described in the method of the inv
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/note- "cloning polylinker prior to a promoterless
chloramphicol acetyl transferase gene"
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                                                                                                                                                                                                                                                                                                                                            AAQ83833 is the plasmid pCM-B64, it was used in the production the bacteriophage T7 RNA polymerase mutant promoter phi 10 -8T to C substitution (claimed).
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to C) prior to a promoterless chloramphicol
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                                           AAQ83834 is the plasmid pCM-P1208, it was used the bacteriophage T7 RNA polymerase mutant pron to G substitution (claimed).
                                                                                                                                                                       New mutant form of T7 RNA polymerase - able to utilise a wide range of T7 promoter sequences having mutations that inactivate recognition by wild type enzyme
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                                        (GEOR-) GEORGIA TECH RES CORP
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/note= "inactive T7
A) prior to
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96.5%;
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10; circular; cycli
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No. 4.9e-78;
                                                                                                                                                     promoter phi 10 mutant (-7C a promoterless chloramphicol
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Columns 111-118; 79pp;
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                                               Promoter
                                                                                                                        Plasmid pCM-T221
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413; Conserv
                                                                                             d pCM-T221;
-6A to G; T
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198..220
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         "intermediate T7 promoter phi 10 mutant (-6A to G) prior to a promoterless chloramphicol acetyl transferase gene"
                                                                                                                                                                                 5110
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Pred. No. 4.9e-78;
0; Mismatches 15;
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Plasmid 04-OCT-1995

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                                    aagcagaaggccatcctgacggatggcctttttgcgtttctacaaactcttcctgtcgtc
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promoters; circular; cyclic; ds.
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T7 RNA polymerase mutant
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Pred. No. 4.9e-78;
0; Mismatches 15;
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                                                                                                                                                                                                                                                                                                                                                                                                           the production promoters.
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                                                                                                                                                                                                                                                                                                   New mutant form of T7 RNA polymerase - able to utilise a range of T7 promoter sequences having mutations that inac recognition by wild type enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mutant promoters;
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                                                                                                          tgaaacgccgtagcgccgatggtagtgtggggtctccccatgcgagagtagggaactgcc
                                                                                                                                                          agagtcgacctgcaggcatgcaagcttgggtcccacctgaccccatgccgaactcagaag
       aagcaacggcccggagggtggcgggcaggacgcccgccataaactgccaggcatcaaatt
                                                                                 aggcatcaaataaaacgaaaggctcagtcgaaagactgggcctttcgttttatctgttgt
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                                            ttgtcggtgaacgctctcctgagtaggacaaatccgccgggagcggatttgaacgttgcg
                                                                       aggcatcaaataaaacgaaaggctcagtcgaaagactgggcctttcgtttatctgttgt
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                                                                                                                                                                                       Conservative
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chloramphicol acetyl transferase
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circular; cyclic;
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T7 promoter phi
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                                                                                                                                                                                                                                                                               Sequence 5110 BP; 1305 A; 1288 C; 1288 G;
                                                                                                                                                                                                                                                                                                                                          AAQ83827 is the plasmid pCM-T297, it was used in the the bacteriophage T7 RNA polymerase mutant promoter p
                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Columns 57-62; 79pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                              New mutant form of T7 RNA polymerase - able to utilise range of T7 promoter sequences having mutations that ir recognition by wild type enzyme
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                                                                                                                                                                       Local Similarity
nes 413; Conserv
                                                                                                                                                                                                                                                                                                                         bacteriophage T7 RNA polymerase mutant substitution (claimed).
agagtcgacctgcaggcatgcaagcttgggtcccacctgaccccatgccgaactcagaag
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/note= "inactive T7 promoter phi 10 mutant (-11G
T) prior to a promoterless chloramphicol
acetyl transferase gene"
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T7 promoter phi 10; circular; cyclic;
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Pred. No. 4.9e-78;
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AAQ83828 is the plasmid pCM-P1160, i the bacteriophage T7 RNA polymerase to G substitution (claimed).
                                                                             New mutant form of T7 RNA polymerase - range of T7 promoter sequences having \pi recognition by wild type enzyme
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                                                  Disclosure; Columns 63-70; 79pp; English.
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G) prior to a promoterless chloramphicol
acetyl transferase gene"
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                                                                                                                                                                                                                 Plasmid pCM-T270; bacteriophage T7; RNA polymerase; mutant -9C to A; T7 promoter phi 10; circular; cycl
                                                                                                                                                                                                                                               Plasmid pCM-T270
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                                                13-AUG-1993;
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                                                                                                                                                                   Promoter
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                            (GEOR-) GEORGIA TECH RES
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Pred. No. 4.9e-78;
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Best Local Similarity 96.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ83829 is the plasmid pCM-T270, it the bacteriophage T7 RNA polymerase to A substitution (claimed).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Columns 71-76; 79pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New mutant form of T7 RNA polymerase - able to utilise a wide range of T7 promoter sequences having mutations that inactivate recognition by wild type enzyme
                                                                                               Promoter
                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                            05-OCT-1995
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Pred. No. 4.9e-78;
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Search completed: September 12, 2002, 09:52:52 Job time: 686 sec